

SEQUENCE LISTING

Masashi WARASHINA
Tomoko WARASHINA

120 Nucleic acid enzymes acquiring an activity for cleaving a target RNA by recognizing another molecule

130

140

-141

150 JP 2000-313320

151 2000-10-13

160 17

-170 PatentIn Ver. 2.0

210 - 1

.211 - 32

+212 + RNA

~213 Artificial Sequence

 $\pm 220^{\circ}$

-223 Description of Artificial Sequence: maxizyme-constituting RNA mole

cule

·3400> 1

ggueruggee ugaugagagu gaugageueu ue 32

 $\pm 210 - 2$

[211] [27]

- 2121- RNA

-213 Artificial Sequence

220

 ± 223 Description of Artificial Sequence: maxizyme-constituting RNA mole cule

[400 - 2]

gucugacugu ucauegaaac eggguee

27

-210 - 3

211 - 33

212 - RNA

-213 - Artificial Sequence

· 220 ·

 $\pm 223 \pm \text{Description}$ of Artificial Sequence: maxizyme-constituting RNA molecule

-1400. -3

ggueeuggee ugaugagagu uauugauggu eag

33

(210) - 4211 + 29 $212 \leq RNA$ 213 Artificial Sequence 220 :223 Description of Artificial Sequence: maxizyme-constituting RNA mole cule -400 - 429 gaagggeuue uuucaucgaa accggguee +210 - 5211 - 88212 RNA 213 Artificial Sequence 220 223 Description of Artificial Sequence: $tRNA^{v_{al}}$ promoter sequence 400 - 5accguugguu uccguagugu agugguuauc acguucgccu aacacgcgaa aggucccegg 60 nucgaaaccg ggcacuacaa aaaccaac 88 210 6 211 33

212 RNA

2135 Artificial Sequence

220 223 Description of Artificial Sequence: ribozyme 220. 223 n is a, c, g or u. 400 6 33 nnnnncugau gaggeegaaa ggeegaaann nnn 2101 - 72111 24 2121 RNA 2131 Artificial Sequence 220 223] Description of Artificial Sequence: left side sequence of maxizyme 4001 7 egaugaecug augagegaaa egge 24 $\pm .2101 - 8$ [211] 24 +12121 RNA E213 Artificial Sequence 2201

4/8

223] Description of Artificial Sequence: right side sequence

of maxizyme

| (400) | 8 | |
|--------|---|----|
| egggg | cugau gagegaaaeg uuce | 24 |
| | | |
| -[210] | 9 | |
| 211 | 13 | |
| - 2121 | RNA | |
| - 2130 | Artificial Sequence | |
| | | |
| 2201 | | |
| - 2231 | Description of Artificial Sequence: substrate | |
| | | |
| 400 | 9 | |
| geegue | eguca ueg | 13 |
| | | |
| (210) | 10 | |
| - 211 | 11 | |
| - 212 | RNA | |
| - 213 | Artificial Sequence | |
| | | |
| [220] | | |
| - 223 | Description of Artificial Sequence: substrate | |
| | | |
| 400 | 10 | |
| geegue | recec g | 11 |
| | | |
| . 210 | 11 | |

211 - 15

.212. RNA

- 213 Artificial Sequence
- 2201
- 223: Description of Artificial Sequence: substrate
- 400 11

ggaacguegu egueg

15

- 2101-12
- 211 40
- 212 RNA
- -213 Artificial Sequence
- 220
- ± 223 Description of Artificial Sequence: wild type ribozyme
- $\pm 400 \pm 12$

ggueeuggee ugaugaggee gaaaggeega aaceggguee

40

- $\pm 210 \leq 13$
- -211 19
- $\pm 212 \pm RNA$
- ·213 · Artificial Sequence
- × 220 ×
- ± 223 Description of Artificial Sequence: part of bc1-2 mRNA as a substrate

1400 - 13

- 210 14
- 2111 25
- + 2121 RNA
- +213 Artificial Sequence
- $+220^{\circ}$
- +223% Description of Artificial Sequence: part of HIV tat mRNA
- -400 14

gaagagcuca ucagaacagu cagac

25

- 210 15
- -211 28
- + 212 RNA
- -213 Artificial Sequence
- -220
- +223 Description of Artificial Sequence: part of BCR-ABL mRNA
- -1400 15
- cugaccauca auaaggaaga agcecuuc

28

- $\pm 1210 \pm 16$
- -1211 20
- 1212 RNA
- E213: Artificial Sequence

| | 0 | 9 | 0 | ١, |
|--|---|---|---|----|

- [223] Description of Artificial Sequence: part of normal ABL mRNA
- 400. 16

uuau uggaa gaageeeuue

20

- 2101 17
- +2111 138
- + 2121 RNA
- -213: Artificial Sequence

220

 ${<}223{>}$ Description of Artificial Sequence: tRNA $^{\rm Val}$ T-MzL

 $\pm 400 \pm 17$

aceguugguu uceguagugu agugguuauc acguucgccu aacacgcgaa agguccecgg 60 uucgaaaccg ggcacuacaa aaaccaacuu ugucugacug uucaucgaaa ceggguecgg 120 uaceeeggau aucuuuuu